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(FILE 'HOME' ENTERED AT 13:15:51 ON 15 APR 2001)

FILE 'BIOSIS, CAPLUS, EMBASE, MEDLINE, SCISEARCH' ENTERED AT 13:16:36 ON 15 APR 2001

```
L1
              3 S FELL PROTEIN
L2
            388 S HYALURONATE (P) (BINDING (W) PROTEIN)
L3
            584 S L2 OR HABP
              O S L3 (P) FELL
L4
              0 S L3 AND L1
L5
L6
             30 S L3 (P) CD44
L7
              0 S L3 (P) ((CD44 (S) PRECURSOR))
Γŝ
              0 S L3 (P) ((CD44 (W)LIKE))
```

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Version:
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 	Туре	٦ +	Hits	Search Text	DBs	Time Stamp	Comment s	Error Definition	E L
	BRS	L	0	FELL adj protein	USPA T; US-P GPUB ; JPO; JPO; DERW	2001/04/1 5 12:31			0
ļ	BRS	L7	23	CD44 same precursor	USPA T; US-P GPUB ; JPO; DERW ENT	2001/04/1 5 12:38			0
	BRS	L19	7	L13 same CD44	USPA T; US-P GPUB ; HPO; JPO; DERW	2001/04/1 5 12:41			0
j	BRS	L13	16	hyaluronate with (binding adj protein)	USPA T; US-P GPUB; HPO; JPO; DERW	2001/04/1 5 12:50			0

Hits S6		ear	Search Text	DBs	Time Stamp	Comment s	Error Definition	Er ro rs
125 2	2 2 6	L13 or HABP		USPA T; US-P GPUB ; JPO; JPO; DERW	2001/04/15 12:51		·	0





National Center for Biotechnology Information

National Library of Medicine

National Institutes of Health

PubMed

Entrez

BLAST OMIMO Taxonomy

Structure

Search GenBank for

Go

SITE MAP

About NCBI general and contact information

GenBank sequence submission support and software

Molecular databases sequences, structures and taxonomy

Literature databases PubMed and **OMIMO**

Genomic biology the human genome, whole genomes and related resources

Tools for data mining

Research at **NCBI** people, projects and seminars

Education

What does NCBI do?

Established in 1988 as a national resource for molecular biology information, NCBI creates public databases, conducts research in computational biology, develops software tools for analyzing genome data, and disseminates biomedical information - all for the better understanding of molecular processes affecting human health and disease.

Draft Human Genome

Explore human genome resources or browse the human genome sequence using the Map Viewer.

Integrated STS reports

A:drives



UniSTS presents marker information collected from public resources including GenBank, RHdb, GDB,

and various maps. Zero in on primer and mapping data, e-PCR results, and Map Viewer and LocusLink crossreferences. More ...

NCBI in the News

The draft sequences of the human genome were compared in an article by Aach et al. (Nature, Feb.

Hot Spots

- Cancer aenome anatomy project
- Clusters of orthologous groups
- ▶ Coffee Break
- Electronic **PCR**
- ▶ Gene expression omnibus
- Genes and disease
- ▶ Human genome resources
- Human/mouse homology maps
- LocusLink
- Malaria genetics & genomics
- ORF finder
- Reference sequence project

teaching resources and on-line tutorials

FTP site download data and software

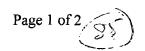
15), noting overall similarities but differences in details. Use of RefSeq, NCBI's manually curated database of mRNA sequences, is credited with providing additional gene annotation for the public sequence.

Disclaimer Privacy statement

Revised April 4, 2001

Retrovirus resources

- ▶ Serial analysis of gene expression
- ▶ Trace archive
- ▶ UniGene
- ▶ VecScreen



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Credits

Mirror sites

<u>Jobs</u>

About GeneCards

Disease genes

How to Mirror

Guiding the user

Data Sources

Data Extraction

Usage statistics

Citing this resource

Publications

What's New

Version: 2.19 Release: Jan 31, 2001 Entries: 18,583

Approved*:11,980

About Bioinformatics
Data Mining in
Biology
Web Usability
Science on the Web

Your Feedback

GeneCards: human genes, maps, proteins and diseases

GeneCards is a database of human genes, their products and their involvement in diseases. It offers concise information about the functions of all human genes that have an approved symbol, as well as selected others [gene listing].

GeneCards now also supports searching <u>UDB</u> (The Unified Database for Human Genome Mapping). <u>Read more about UDB</u>.

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[Quick Start] [Guided Tour] [More search examples]

• Search/Display GeneCards by keyword(s)

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For example, you can display the GeneCard for the (case-sensitive) symbol BRCA1.

or search GeneCards for the keyword(s):

- o p53
- o apolipoprot*AND (hyper* OR Alzheim*)
- o U85267 GenBank accession No.
- o Hs.1288 UniGene cluster
- o ATCC: 106253, image: 303124 clone identifier
- o chromosome: 22, locus: 20p*,locus: 7p13
- Search UDB integrated map

specify chromosome: 1 Go For example, a map region in chromosome 19.

- Search UDB by gene/marker name

 For example, Information about the mapped marker D17S1843.
- View estimated boundaries (in Megabases) of cytogenetic bands

specify chromosome: 1 GG

What's special about GeneCards?

The <u>information presented</u> here has been <u>automatically extracted</u> from various <u>resources</u> by scripts developed in <u>our group</u>. GeneCards is

particulary useful for people who wish to find information about genes of interest in the context of <u>functional genomics</u> and proteomics.

This resource also features a new type of <u>navigation support</u> system that **guides** its <u>users</u> to the information. Important parts of this guidance system are the **spell corrector**, and the automatically generated tips for query reformulation.

Developed at the Crown Human Genome Center & Bioinformatics Unit, at the Weizmann Institute of Science

Credits:

Michael Rebhan, Avital Adato, Vered Chalifa-Caspi, Inga Peter, Jaime Prilusky, Michael Ronen, Hershel Safer, Marilyn Safran, Shai Shen-Orr, Liora Yaar, Doron Lancet

Comments to: cards@bioinfo.weizmann.ac.il

* Entries with HUGO-approved symbols. <u>Back to top</u>